SEQUENCE LISTING

<110> WEI, Ming-Hui

<120> ISOLATED HUMAN TRANSPORTER PROTEINS,
    NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
    AND USES THEREOF

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14

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15

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16

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Val Gly Leu Phe Gly Thr Met Ile Met Gly Ala Ile Cys Thr His Cys
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Met His Met Leu Val Asn Cys Ser His Glu Leu Cys Arg Arg Phe Gln
    50                  55                  60
Gln Pro Ser Leu Asp Phe Ser Glu Val Ala Tyr Cys Ser Phe Glu Ser
65                  70                  75                  80
Gly Pro Leu Gly Leu Arg Arg Tyr Ser Met Leu Ala Arg Arg Ile Val
                85                  90                  95
Thr Thr Phe Leu Phe Ile Thr Gln Ile Gly Phe Cys Cys Val Tyr Phe
            100                 105                 110
Leu Phe Val Ala Leu Asn Ile Lys Asp Val Met Asp His Tyr Tyr Lys
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Met Pro Val Gln Ile Tyr Leu Leu Ile Met Leu Gly Pro Met Ile Leu
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Ala Ala Leu Leu Thr Val Ala Gly Leu Ala Ile Thr Phe Ser Tyr Met
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Leu Val Asp Leu Pro Asp Val His Thr Val Lys Pro Val Ala Thr Trp
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Ala Thr Leu Pro Leu Tyr Phe Gly Thr Ala Ile Tyr Ala Phe Glu Gly
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His Val Glu Gly Ser Ile Thr Leu Asn Leu Pro Gln Gly Asp Thr Leu
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Ser Gln Leu Val Arg Ile Ser Met Ala Val Ala Ile Phe Leu Ser Tyr
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Arg Val Val Leu Val Thr Phe Thr Phe Leu Leu Ala Thr Cys Ile Pro
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17

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        35                  40                  45
Met His Met Leu Val Asn Cys Ser His Glu Leu Cys Arg Arg Phe Gln
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Gln Pro Ser Leu Asp Phe Ser Glu Val Ala Tyr Cys Ser Phe Glu Ser
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18

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Ala Gln Ile Leu

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Cys Leu Ile Cys Leu Tyr Cys Met Arg Gln Val Val Phe Ala Ala His
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Phe Val Cys Asn Arg Asn Gly Arg Asp Leu Ile Asp Tyr Ala Asn Ile
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          165   170   175
Ala Val Ala Val Val Leu Phe Phe Phe Leu Ser Asp Leu Arg Pro Ile
          180   185   190
Ser Ser Leu Pro Trp Phe Gly Lys Ala Thr Asp Leu Pro Leu Phe Phe
          195   200   205
Gly Thr Val Met Phe Ala Phe Glu Gly Val Ala Val Ile Met Pro Ile
        210   215   220
Glu Asn Arg Met Gln Ser Pro His Ala Phe Ile Ser Trp Asn Gly Val
225   230   235   240
Leu Asn Ser Ser Cys Leu Val Val Leu Ala Ile Phe Ser Val Thr Gly
          245   250   255
Phe Tyr Gly Tyr Leu Ser Leu Gly Asn Asp Val Lys Asp Thr Ala Thr
          260   265   270
Leu Asn Leu Pro Met Thr Pro Phe Tyr Gln Thr Ile Lys Leu Met Phe
        275   280   285
Val Ala Cys Ile Met Ile Ser Tyr Pro Leu Gln Phe Tyr Val Pro Met
        290   295   300
Glu Arg Ile Glu Lys Trp Ile Thr Arg Lys Ile Pro Val Asp Lys Gln

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305                    310                    315                    320
Thr Leu Tyr Ile Tyr Ile Ala Arg Tyr Ser Gly Val Ile Leu Thr Cys
                    325                    330                    335
Ala Ile Ala Glu Leu Ile Pro His Leu Ala Leu Phe Ile Ser Leu Ile
                340                    345                    350
Gly Ala Phe Ser Gly Ala Ser Met Ala Leu Leu Phe Pro Pro Cys Ile
            355                    360                    365
Glu Leu Leu Thr Ser Tyr Ala Lys Asn Glu Leu Ser Thr Gly Leu Trp
        370                    375                    380
Ile Lys Asn Ile Val Leu Leu Thr Phe Ala Phe Ile Gly Phe Thr Thr
385                    390                    395                    400
Gly Thr Tyr
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<400> 7
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Thr Asp Val Ser Pro Glu Glu Ser Pro Ser Glu Gly Leu Asn Asn Leu
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Ser Ser Pro Gly Ser Tyr Gln Arg Phe Gly Gln Ser Asn Ser Thr Thr
        35                   40                   45
Trp Phe Gln Thr Leu Ile His Leu Leu Lys Gly Asn Ile Gly Thr Gly
    50                   55                   60
Leu Leu Gly Leu Pro Leu Ala Val Lys Asn Ala Gly Ile Val Met Gly
65                   70                   75                   80
Pro Ile Ser Leu Leu Ile Ile Gly Ile Val Ala Val His Cys Met Gly
                85                   90                   95
Ile Leu Val Lys Cys Ala His His Phe Cys Arg Arg Leu Asn Lys Ser
            100                  105                  110
Phe Val Asp Tyr Gly Asp Thr Val Met Tyr Gly Leu Glu Ser Ser Pro
        115                  120                  125
Cys Ser Trp Leu Arg Asn His Ala His Trp Gly Arg Arg Val Val Asp
    130                  135                  140
Phe Phe Leu Ile Val Thr Gln Leu Gly Phe Cys Cys Val Tyr Phe Val
145                  150                  155                  160
Phe Leu Ala Asp Asn Phe Lys Gln Val Ile Glu Ala Ala Asn Gly Thr
                165                  170                  175
Thr Asn Asn Cys His Asn Asn Glu Thr Val Ile Leu Thr Pro Thr Met
            180                  185                  190
Asp Ser Arg Leu Tyr Met Leu Ser Phe Leu Pro Phe Leu Val Leu Leu
        195                  200                  205
Val Phe Ile Arg Asn Leu Arg Ala Leu Ser Ile Phe Ser Leu Leu Ala
    210                  215                  220
Asn Ile Thr Met Leu Val Ser Leu Val Met Ile Tyr Gln Phe Ile Val
225                  230                  235                  240
Gln Arg Ile Pro Asp Pro Ser His Leu Pro Leu Val Ala Pro Trp Lys
                245                  250                  255
Thr Tyr Pro Leu Phe Phe Gly Thr Ala Ile Phe Ser Phe Glu Gly Ile
            260                  265                  270
Gly Met Val Leu Pro Leu Glu Asn Lys Met Lys Asp Pro Arg Lys Phe
        275                  280                  285
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20

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Pro Leu Ile Leu Tyr Leu Gly Met Val Ile Val Thr Ile Leu Tyr Ile
    290                 295             300
Ser Leu Gly Cys Leu Gly Tyr Leu Gln Phe Gly Ala Asn Ile Gln Gly
305             310                 315                     320
Ser Ile Thr Leu Asn Leu Pro Asn Cys Trp Leu Tyr Gln Ser Val Glu
                325                 330                     335
Leu Leu Tyr Leu Gly Gly Ile Cys Leu Thr Tyr Pro Leu Gln Phe Tyr
            340                 345                 350
Val Ser Ala Lys Ile Ile Val Pro Val Ile Val Ser Trp Val Cys Lys
            355                 360                 365
Cys Cys Thr Leu Met Val Asp Leu Gly Ile Gly Ser Ala Met Leu Cys
    370                 375                 380
Lys Thr Cys Ile Leu Ala Ile Leu Ile Pro Arg Leu Asp Leu Val Ile
385                 390                 395                     400
Ser Leu Val Gly Ser Val Ser Ser Ser Ala Leu Ala Leu Ile Ile Pro
                405                 410                 415
Pro Leu Leu Glu Val Thr Thr Phe Tyr Ser Glu Gly Met Ser Pro Leu
            420                 425                 430
Thr Ile Phe Lys Asp Ala Leu Ile Ser Ile Leu Gly Phe Val Gly Phe
            435                 440                 445
Val Val Gly Thr Tyr Glu Ala Leu Tyr Glu Leu Ile Gln Pro Ser Asn
    450                 455                 460
Ala Pro Ile Phe Ile Asn Ser Thr Cys Ala Phe Ile
465                 470                 475
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